

Appendix A15. AD Model Builder code for the instantaneous rates catch/release model (IRCR).

```
//--><>--><>--><>--><>--><>--><>--><>--><>--><>--><>--><>--><>--> //  
// Jiang et. al (2007) Age-independent instantaneous rates model for catch and release //  
//  
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// Version 1.2 //  
//--><>--><>--><>--><>--><>--><>--><>--><>--><>--><>--><>--><>--><>-->
```

DATA_SECTION

```
// Starting and ending year of the release year  
init_int styrR;  
init_int endyrR;  
  
//Starting and ending year of recovery years  
init_int styr;  
init_int endyr;  
  
//Total Releases by Year  
init_vector N(styrR,endyrR);  
  
//Recapture Matrix for harvest fish  
init_imatrix rh(styrR,endyrR,styr,endyr);  
  
//Recapture Matrix for releases fish  
init_imatrix rr(styrR,endyrR,styr,endyr);  
  
//----Reporting Rate for harvested fish-----  
init_number lh;  
  
//---Initial probability of tag shedding and tag-induced mortality for harvested fish--  
init_number phih;  
  
//---Reporting Rate for released fish-----  
init_number lr;  
  
//---Initial probability of tag shedding and tag-induced mortality for released fish--  
init_number phir;  
  
//Hooking Mortality  
init_number h;  
  
//Number of Natural Mortality Periods and Beginning Years  
init_int mp;  
init_ivector mp_int(1,mp);  
int pp;  
  
//Number of Fishing Mortality Periods and Beginning Years  
init_int fp;  
init_ivector fp_int(1,fp);  
int qq;  
  
//Number of Tag Mortality Periods  
init_int fap;  
init_ivector fap_int(1,fap);  
int ss;  
int tp;
```

LOCAL_CALCS

```
pp=mp+1;  
qq=fp+1;
```

```

ss=fap+1;
tp=mp+fp+fap+(4*(endyr-styr+1));
END_CALCS
matrix sigma(1,tp,1,tp+1);
!! set_covariance_matrix(sigma);
//looping variables
int y;
int t;
int a;
int d;
int cnt;
int total;
int Ntags;
int looper;
int df_r;
int df_h;
int hless;
int rless;

PARAMETER_SECTION
number dodo;
number dodol;
number probs;
number AIC;
number AICc;
number K;
number up_df;
number up_count;
number up_chi;
number up_chat;
number p_chi;
number p_df;
number p_chat;

//-----F estimates-----
init_bounded_vector e_F(1,fp,-30.,1.6,1);
vector F(styr,endyr);
vector fp_yr(1,qq);

//-----M estimates-----
init_bounded_vector e_M(1,mp,-30,1.6,1);
vector M(styr,endyr);
vector mp_yr(1,pp);

//-----Tag Mortality-----
init_bounded_vector e_FA(1,fap,-30.,1.6,1);
vector FA(styr,endyr);
vector fap_yr(1,ss);

//-----Tag Number of Tags-----
vector tags(styrR,endyrR);

//-----Mortality Calculations-----
matrix s(styrR,endyrR,styr,endyr);
matrix u_h(styrR,endyrR,styr,endyr);
matrix u_r(styrR,endyrR,styr,endyr);
vector S_fish(styr,endyr);

//-----Predicted Cell recoveries-----
vector sum_prob_h(styrR,endyrR);
vector sum_prob_r(styrR,endyrR);
matrix s_prob(styrR,endyrR,styr,endyr);
matrix exp_prob_h(styrR,endyrR,styr,endyr);
matrix ll_h(styrR,endyrR,styr,endyr);
matrix exp_prob_r(styrR,endyrR,styr,endyr);
matrix ll_r(styrR,endyrR,styr,endyr);
vector ll_ns(styrR,endyrR);
matrix exp_r_h(styrR,endyrR,styr,endyr);
matrix exp_r_r(styrR,endyrR,styr,endyr);
matrix pool_r(styrR,endyrR,styr,endyr);
matrix pool_h(styrR,endyrR,styr,endyr);

```

```

matrix pool_r_e(styrR,endyrR,styr,endyr);
matrix pool_h_e(styrR,endyrR,styr,endyr);
matrix chi_r(styrR,endyrR,styr,endyr);
matrix chi_h(styrR,endyrR,styr,endyr);
matrix p_chi_r(styrR,endyrR,styr,endyr);
matrix p_chi_h(styrR,endyrR,styr,endyr);
matrix pear_r(styrR,endyrR,styr,endyr);
matrix pear_h(styrR,endyrR,styr,endyr);
vector exp_ns(styrR,endyrR);
vector chi_ns(styrR,endyrR);
vector pear_ns(styrR,endyrR);
sdreport_vector S(styr,endyr);
sdreport_vector FM(styr,endyr);
sdreport_vector FT(styr,endyr);
sdreport_vector NM(styr,endyr);

//-----Likelihood Values-----
number f_tag;
objective_function_value f;

INITIALIZATION_SECTION
e_F -1.6;
e_FA -1.6;
e_M -1.6;

RUNTIME_SECTION
maximum_function_evaluations 100, 500, 5000;
convergence_criteria 1e-5, 1e-7, 1e-16;

PRELIMINARY_CALCS_SECTION
F.initialize();
FA.initialize();
M.initialize();
PROCEDURE_SECTION
calc_number_tags();
calc_M_vector();
calc_F_vector();
calc_FA_vector();
calc_fish_surv();
calc_s();
calc_s_prob();
calc_u_h();
calc_u_r();
calc_exp_prob_h();
calc_exp_prob_r();
calc_LL();
calc_Chisquare();
calc_pooled_cells();
evaluate_the_objective_function();

FUNCTION calc_number_tags
cnt=0;
for(t=styrR;t<=endyrR;t++) {
Ntags=0;
for(y=styr+cnt;y<=endyr;y++) {
Ntags+=rh(t,y)+rr(t,y);
}
tags(t)=Ntags;
cnt+=1;
}

FUNCTION calc_M_vector
for(t=1;t<=mp;t++) {
mp_yr(t)=mp_int(t);
}
mp_yr(pp)=endyr+1;

for(t=styr;t<=endyr;t++) {
for(d=1;d<=mp;d++) {
if(t>=mp_yr(d) && t<mp_yr(d+1)) {
M(t)=mfexp(e_M(d));
}
}
}

```

```

        NM(t)=M(t);
    }
}

FUNCTION calc_F_vector
for(t=1;t<=fp;t++) {
    fp_yr(t)=fp_int(t);
}
fp_yr(qq)=endyr+1;

for(t=styr;t<=endyr;t++) {
    for(d=1;d<=fp;d++) {
        if(t>=fp_yr(d) && t<fp_yr(d+1)) {
            F(t)=mfexp(e_F(d));
            FM(t)=F(t);
        }
    }
}

FUNCTION calc_FA_vector
for(t=1;t<=fap;t++) {
    fap_yr(t)=fap_int(t);
}
fap_yr(ss)=endyr+1;

for(t=styr;t<=endyr;t++) {
    for(d=1;d<=fap;d++) {
        if(t>=fap_yr(d) && t<fap_yr(d+1)) {
            FA(t)=mfexp(e_FA(d));
            FT(t)=FA(t);
        }
    }
}

FUNCTION calc_fish_surv
for (t=styr;t<=endyr;t++) {
    S_fish(t)=mfexp(-1*(F(t)+h*FA(t)+M(t)));
    S(t)=S_fish(t);
}

FUNCTION calc_s
cnt=0;
for(t=styrR;t<=endyrR;t++) {
    for(y=styr+cnt;y<=endyr;y++) {
        if(t==y){s(t,y)=1;}
        if(t!=y){s(t,y)=mfexp(-F(y)-FA(y)-M(y));}
    }
    cnt+=1;
}

FUNCTION calc_u_h
cnt=0;
for(t=styrR;t<=endyrR;t++) {
    for(y=styr+cnt;y<=endyr;y++) {
        u_h(t,y)=(F(y)/(F(y)+FA(y)+M(y)))*(1-mfexp(-F(y)-FA(y)-M(y)));
    }
    cnt+=1;
}

FUNCTION calc_u_r
cnt=0;
for (t=styrR;t<=endyrR;t++) {
    for (y=styr+cnt;y<=endyr;y++) {
        u_r(t,y)=(FA(y)/(F(y)+FA(y)+M(y)))*(1-mfexp(-F(y)-FA(y)-M(y)));
    }
    cnt+=1;
}

FUNCTION calc_s_prob
cnt=0;

```

```

for(t=styrR;t<=endyrR;t++) {
    looper=0;
    for(y=styr+cnt;y<=endyr;y++) {
        probs=1;
        for(a=y-looper;a<=y;a++) {
            probs=probs*s(t,a);
        }
        s_prob(t,y)=probs;
        looper+=1;
    }
    cnt+=1;
}

FUNCTION calc_exp_prob_h
cnt=0;
for(t=styrR;t<=endyrR;t++) {
    dodo=0;
    for(y=styr+cnt;y<=endyr;y++) {
        exp_prob_h(t,y)=lh*phih*s_prob(t,y)*u_h(t,y);
        dodo+=exp_prob_h(t,y);
    }
    sum_prob_h(t)=dodo;
    cnt+=1;
}

FUNCTION calc_exp_prob_r
cnt=0;
for(t=styrR;t<=endyrR;t++) {
    dodo=0;
    for(y=styr+cnt;y<=endyr;y++) {
        exp_prob_r(t,y)=lr*phir*s_prob(t,y)*u_r(t,y);
        dodo+=exp_prob_r(t,y);
    }
    sum_prob_r(t)=dodo;
    cnt+=1;
}

FUNCTION calc_LL
cnt=0;
for(t=styrR;t<=endyrR;t++) {
    for(y=styr+cnt;y<=endyr;y++) {
        ll_h(t,y)=0;
        ll_r(t,y)=0;
        if(rh(t,y)!=0) {
            ll_h(t,y)=rh(t,y)*log(exp_prob_h(t,y));
        }
        if(rr(t,y)!=0) {
            ll_r(t,y)=rr(t,y)*log(exp_prob_r(t,y));
        }
    }
    cnt+=1;
}
for (t=styrR;t<=endyrR;t++) {
    ll_ns(t)=(N(t)-tags(t))*log(1-(sum_prob_h(t)+sum_prob_r(t)));
}

FUNCTION evaluate_the_objective_function
f_tag=0;
cnt=0;
for(t=styrR;t<=endyrR;t++) {
    for(y=styr+cnt;y<=endyr;y++) {
        f_tag+=ll_h(t,y)+ll_r(t,y);
    }
    cnt+=1;
}

for(t=styrR;t<=endyrR;t++) {
    f_tag+=ll_ns(t);
}
f=f_tag*-1.;

```

```

FUNCTION calc_Chisquare
cnt=0;
up_count=0;
for(t=styrR;t<=endyrR;t++) {
    for(y=styr+cnt;y<=endyr;y++) {
        up_count+=1;
    }
    cnt+=1;
}

cnt=0;
for(t=styrR;t<=endyrR;t++) {
    for(y=styr+cnt;y<=endyr;y++) {
        exp_r_r(t,y)=exp_prob_r(t,y)*N(t);
        exp_r_h(t,y)=exp_prob_h(t,y)*N(t);
    }
    cnt+=1;
}

cnt=0;
for(t=styrR;t<=endyrR;t++) {
    for(y=styr+cnt;y<=endyr;y++) {
        chi_r(t,y)=square(rr(t,y)-exp_r_r(t,y))/exp_r_r(t,y);
        chi_h(t,y)=square(rh(t,y)-exp_r_h(t,y))/exp_r_h(t,y);
        pear_r(t,y)=(rr(t,y)-exp_r_r(t,y))/sqrt(exp_r_r(t,y));
        pear_h(t,y)=(rh(t,y)-exp_r_h(t,y))/sqrt(exp_r_h(t,y));
    }
    cnt+=1;
}
for (t=styrR;t<=endyrR){
    exp_ns(t)=N(t)*(1-(sum_prob_h(t)+sum_prob_r(t)));
}

//Not seen chi
for(t=styrR;t<=endyrR;t++) {
    chi_ns(t)=0;
    chi_ns(t)=square((N(t)-tags(t))-exp_ns(t))/exp_ns(t);
    pear_ns(t)=((N(t)-tags(t))-exp_ns(t))/sqrt(exp_ns(t));
}

//total chi square
up_chi=sum(chi_r)+sum(chi_h)+sum(chi_ns);
K=fap+mp+fp;
up_df=up_count*2-K;
up_chat=up_chi/up_df;
AIC=-1.*2*f_tag+2*K;
AICc=AIC+(2*K*(K+1))/(sum(N)-K-1);

FUNCTION calc_pooled_cells
// Pool harvested cells
cnt=0;
for(t=styrR;t<=endyrR;t++) {
    for(y=styr+cnt;y<=endyr;y++) {
        pool_h_e(t,y)=0;
        pool_h(t,y)=0;
        pool_h_e(t,y)=exp_r_h(t,y);
        pool_h(t,y)=rh(t,y);
    }
    cnt+=1;
}
cnt=0;
hless=0;
for(t=styrR;t<=endyrR;t++) {
    for(y=endyr;y>=styr+cnt;y--) {
        if(pool_h_e(t,y)>=1) {
            pool_h(t,y)=pool_h(t,y);
            pool_h_e(t,y)=pool_h_e(t,y);
        }
        if(pool_h_e(t,y)>=0 && pool_h_e(t,y)<1) {
            if(y!=styr+cnt)
            {

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        hless+=1;
        pool_h_e(t,y-1)=pool_h_e(t,y-1)+pool_h_e(t,y);
        pool_n(t,y-1)=pool_h(t,y-1)+pool_h(t,y);
        pool_h(t,y)=0;
        pool_h_e(t,y)=0;
    }
    if (y==styr+cnt) break;
}
}//for
cnt+=1;
}//for

// Pool released cells
cnt=0;
for(t=styrR;t<=endyrR;t++) {
    for(y=styr+cnt;y<=endyr;y++) {
        pool_r_e(t,y)=0;
        pool_r(t,y)=0;
        pool_r_e(t,y)=exp_r_r(t,y);
        pool_r(t,y)=rr(t,y);
    }
    cnt+=1;
}
cnt=0;
rless=0;
for(t=styrR;t<=endyrR;t++) {
    for(y=endyr;y>=styr+cnt;y--) {
        if(pool_r_e(t,y)>=1) {
            pool_r(t,y)=pool_r(t,y);
            pool_r_e(t,y)=pool_r_e(t,y);
        }
        if(pool_r_e(t,y)>=0 && pool_r_e(t,y)<1) {
            if (y!=styr+cnt) {
                rless+=1;
                pool_r_e(t,y-1)=pool_r_e(t,y-1)+pool_r_e(t,y);
                pool_r(t,y-1)=pool_r(t,y-1)+pool_r(t,y);
                pool_r(t,y)=0;
                pool_r_e(t,y)=0;
            }
            if (y==styr+cnt) break;
        }
    }
    cnt+=1;
}
p_df=up_count*2-hless-rless-K;

//Pooled Chi-square
cnt=0;
for(t=styrR;t<=endyrR;t++) {
    for(y=styr+cnt;y<=endyr;y++) {
        p_chi_h(t,y)=0;
        p_chi_r(t,y)=0;
        if(pool_h_e(t,y)!=0) {
            p_chi_h(t,y)=square(pool_h(t,y)-pool_h_e(t,y))/pool_h_e(t,y);
        }
        if(pool_r_e(t,y)!=0) {
            p_chi_r(t,y)=square(pool_r(t,y)-pool_r_e(t,y))/pool_r_e(t,y);
        }
    }
    cnt+=1;
}
p_chi=sum(p_chi_h)+sum(p_chi_r)+sum(chi_ns);
p_chat=p_chi/p_df;

REPORT_SECTION
report<<"Log-L<<"<<"\t"<<"K"<<"\t"<<"AIC"<<"\t"<<"AICc"<<"\t"<<"Eff. Sample
Size"<<endl;
report<<f_tag<<"<<"\t"<<K<<"\t"<<AIC<<"\t"<<AICc<<"\t"<<sum(N)<<endl;
report<<"<<endl;
report<<"<<endl;
report<<"*****Model Statistics*****"<<endl;

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report<<"Unpooled Chi-square      "<<"    "<<up_chi<<endl;
report<<"Unpooled df           "<<"    "<<up_df<<endl;
report<<"Unpooled c-hat         "<<"    "<<up_chat<<endl;
report<<"Pooled Chi-square      "<<"    "<<p_chi<<endl;
report<<"Pooled df             "<<"    "<<p_df<<endl;
report<<"Pooled c-hat          "<<"    "<<p_chat<<endl;
report <<"*****Observed and Calculated Data*****"<<endl;
report << "Obs Recoveries of harvest fish "<< endl;
report<<rh<<endl;
report <<" " <<endl;
report << "Obs Recoveries of release fish "<< endl;
report<<rr<<endl;
report <<" " <<endl;

report << "Total Released "<< endl;
report<<N<<endl;
report <<" " <<endl;

report <<"Total Recovered Tags"<<endl;
report <<tags<<endl;
report <<" " <<endl;

report << "s matrix" << endl;
report <<s<<endl;
report<<" " <<endl;

report << "S_prob matrix" << endl;
report <<s_prob<<endl;
report <<" " <<endl;

report << "Exploitation Rate of harvested fish" << endl;
report <<uh<<endl;
report <<" " <<endl;

report << "Exploitation Rate of released fish" << endl;
report <<ur<<endl;
report <<" " <<endl;

report <<"Expected Probability of harvested fish"<<endl;
report<<exp_prob_h<<endl;
report <<" " <<endl;

report <<"Expected Probability of released fish"<<endl;
report<<exp_prob_r<<endl;
report <<" " <<endl;

report<<"Not Seen Probability"<<endl;
report<<1-(sum_prob_h+sum_prob_r)<<endl;
report <<" " <<endl;

report <<"Expected Number of harvested fish"<<endl;
report<<exp_r_h<<endl;
report <<" " <<endl;

report <<"Expected Number of released fish"<<endl;
report<<exp_r_r<<endl;
report <<" " <<endl;

report <<"Expected Number of not seen"<<endl;
report<<exp_ns<<endl;
report <<" " <<endl;

report <<"Cell Likelihoods of harvested fish"<<endl;
report<<ll_h<<endl;

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report <<" "<<endl;

report <<"Cell Likelihoods of released fish"<<endl;
report<<ll_r<<endl;
report <<" "<<endl;

report <<"Cell Likelihoods of unseen"<<endl;
report<<ll_ns<<endl;
report <<" "<<endl;

report <<"Unpooled Chi-squares of Harvested Fish"<<endl;
report<<chi_h<<endl;
report <<" "<<endl;

report <<"Unpooled Chi-squares of Released Fish"<<endl;
report<<chi_r<<endl;
report <<" "<<endl;

report <<"Chi-squares of Not Seen"<<endl;
report<<chi_ns<<endl;
report <<" "<<endl;

report <<"Pooled Cells of Harvested Fish"<<endl;
report<<pool_h<<endl;
report <<" "<<endl;

report <<"Pooled Expected Cells of Harvested Fish"<<endl;
report<<pool_h_e<<endl;
report <<" "<<endl;

report <<"Pooled Cells of Released Fish"<<endl;
report<<pool_r<<endl;
report <<" "<<endl;

report <<"Pooled Expected Cells of Harvested Fish"<<endl;
report<<pool_r_e<<endl;
report <<" "<<endl;

report <<"Pooled Chi-squares of Harvested Fish"<<endl;
report<<p_chi_h<<endl;
report <<" "<<endl;

report <<"Pooled Chi-squares of Released Fish"<<endl;
report<<p_chi_r<<endl;
report <<" "<<endl;
report <<"Pearson Residuals for released fish"<<endl;
report<<pear_r<<endl;
report <<" "<<endl;

report <<"Pearson Residuals for harvested fish"<<endl;
report<<pear_h<<endl;
report <<" "<<endl;

report <<"Pearson Residuals for not seen"<<endl;
report<<pear_ns<<endl;
report <<" "<<endl;

FINAL_SECTION
//Output F and sd
ofstream ofs1("F.std");
d=mp+fp+fap+(endyr-styr+1);
for(y=styr;y<=endyr;y++){
    d+=1;
    ofs1<<FM(y)<<"\t"<<sigma(d,1)<<endl;
}
//Output FA and sd
ofstream ofs2("FA.std");
for(y=styr;y<=endyr;y++){
    d+=1;
    ofs2<<FT(y)<<"\t"<<sigma(d,1)<<endl;
}

```

```
//Output M and Sd
ofstream ofs3("M.std");
for(y=styr;y<=endyr;y++) {
    d+=1;
    ofs3<<NM(y)<<"\t"<<sigma(d,1)<<endl;
}
```